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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/024,808

DATE: 01/14/2002
TIME: 18:57:21

Input Set : A:\Seqlist.txt
Output Set: N:\CRF3\01142002\J024808.raw

ENTERED

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4 <110> APPLICANT: MEEK, THOMAS
5      PATEL, MEHUL
6      THRALL, SARA D.
8 <120> TITLE OF INVENTION: METHODS OF AND COMPOUNDS FOR MODULATING
9      THE ACTIVITY OF BACTERIAL FabG
12 <130> FILE REFERENCE: P51217
C--> 14 <140> CURRENT APPLICATION NUMBER: US/10/024,808
15 <141> CURRENT FILING DATE: 2001-12-19
17 <150> PRIOR APPLICATION NUMBER: 60/259,595
18 <151> PRIOR FILING DATE: 2000-01-03
20 <160> NUMBER OF SEQ ID NOS: 2
22 <170> SOFTWARE: FastSEQ for Windows Version 4.0
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 243
26 <212> TYPE: PRT
27 <213> ORGANISM: Streptococcus pneumoniae
29 <400> SEQUENCE: 1
30 Met Lys Leu Glu His Lys Asn Ile Phe Ile Thr Gly Ser Ser Arg Gly
31   1           5           10          15
32 Ile Gly Leu Ala Ile Ala His Lys Phe Ala Gln Ala Gly Ala Asn Ile
33   20          25          30
34 Val Leu Asn Ser Arg Gly Ala Ile Ser Glu Glu Leu Leu Ala Glu Phe
35   35          40          45
36 Ser Asn Tyr Gly Ile Lys Val Val Pro Ile Ser Gly Asp Val Ser Asp
37   50          55          60
38 Phe Ala Asp Ala Lys Arg Met Ile Asp Gln Ala Ile Ala Glu Leu Gly
39   65          70          75          80
40 Ser Val Asp Val Leu Val Asn Asn Ala Gly Ile Thr Gln Asp Thr Leu
41   85          90          95
42 Met Leu Lys Met Thr Glu Ala Asp Phe Glu Lys Val Leu Lys Val Asn
43   100         105         110
44 Leu Thr Gly Ala Phe Asn Met Thr Gln Ser Val Leu Lys Pro Met Met
45   115         120         125
46 Lys Ala Arg Glu Gly Ala Ile Ile Asn Met Ser Ser Val Val Gly Leu
47   130         135         140
48 Met Gly Asn Ile Gly Gln Ala Asn Tyr Ala Ala Ser Lys Ala Gly Leu
49   145         150         155         160
50 Ile Gly Phe Thr Lys Ser Val Ala Arg Glu Val Ala Ser Arg Asn Ile
51   165         170         175
52 Arg Val Asn Val Ile Ala Pro Gly Met Ile Glu Ser Asp Met Thr Ala
53   180         185         190
54 Ile Leu Ser Asp Lys Ile Lys Glu Ala Thr Leu Ala Gln Ile Pro Met
55   195         200         205
56 Lys Glu Phe Gly Gln Ala Glu Gln Val Ala Asp Leu Thr Val Phe Leu
57   210         215         220
58 Ala Gly Gln Asp Tyr Leu Thr Gly Gln Val Ile Ala Ile Asp Gly Gly
59   225         230         235         240

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60 Leu Ser Met
64 <210> SEQ ID NO: 2
65 <211> LENGTH: 732
66 <212> TYPE: DNA
67 <213> ORGANISM: Streptococcus pneumoniae
69 <400> SEQUENCE: 2
70 atgaaactag aacataaaaa tatcttatt acaggttcga gtcgtggaat tggcttgcc 60
71 atcgcccaca agtttgcctca agcaggagcc aacatgtct taaacagtct tggggcaatc 120
72 tcagaagaat tgctcgctga gtttcaaac tatggatca aggtggttcc catttcagga 180
73 gatgtatcag atttgcaga cgctaagcgt atgattgatc aagctattgc agaactgggt 240
74 tcagtagatg tttggtcaa caatgcagg attacccaag atactcttat gctcaagatg 300
75 acagaagcag attttgaaaa agtgctcaag gtcaatctga ctgggtgcctt taatatgaca 360
76 caatcagtct tgaaaccgat gatgaaagcc agagaaggt ctatcattaa tatgtctagt 420
77 gttgttggtt tcatggggaa tattggtcaa gctaactatg ctgcttctaa ggctggctt 480
78 attggcttta ccaagtctgt ggacacgcgag gtcgctagtc ggaatatacg agtcaatgtg 540
79 attgctccag gaatgattga gtctgatatg acagctatct tatcagataa gattaaggaa 600
80 gctacactag ctcagattcc gatgaaagaa tttgggcagg cagagcagg tgcagattt 660
81 acagtattt tagcaggcca agattatcta actggtcaag tgattgccat tgatggtggc 720
82 ttaagtatgt ag 732

VERIFICATION SUMMARY
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L:14 M:270 C: Current Application Number differs, Replaced Current Application Number